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(21) International Application Number: PCT/US91/04676 (22) International Filing Date: 1 July 1991 (01.07.91) (30) Priority data: 547,990 2 July 1990 (02.07.90) US (71) Applicant: THE REGENTS OF THE UNIVERSITY OF CALIFORNIA [US/US]; Office of The President, 300 Lakeside Drive, Oakland, CA 94612-3550 (US). (72) Inventors: TSIEN, Roger, Y. ; 8535 Nottingham Place, La Jolla, CA 92037 (US). TAYLOR, Susan, S. ; 1408 Via Alta, Del Mar, CA 92014 (US). ADAMS, Stephen, R. ; 13562 Comuna Drive, Poway, CA 92064 (US). JI, Ying ; 8155 Cargill Avenue, #26, San Diego, CA 92122 (US).		(74) Agents: CAMPBELL, Cathryn et al.; Pretty, Schroeder, Brueggemann & Clark, 444 South Flower Street, Suite 2000, Los Angeles, CA 90071 (US). (81) Designated States: AT (European patent), BE (European patent), CH (European patent), DE (European patent), DK (European patent), ES (European patent), FR (European patent), GB (European patent), GR (European patent), IT (European patent), JP, LU (European patent), NL (European patent), SE (European patent). Published <i>With international search report.</i> <i>Before the expiration of the time limit for amending the claims and to be republished in the event of the receipt of amendments.</i>
(54) Title: DETECTION OF ANALYTES USING FLUORESCENT ENERGY TRANSFER (57) Abstract <p>The invention provides labeled proteins suitable for determining the presence of cAMP, other second messengers and organic molecules. The proteins are separately labeled with fluorochromes which, when in close spatial proximity, preferably, less than about 6 nm, interact through the transfer of energy from one fluorochrome to the other. A composition of matter $(S_1kA)_{n_1}(S_2kD)_{n_2}$ is provided wherein S_1 and S_2 are two proteins which are associated in one state and substantially disassociated in another, the equilibrium between which is controlled by the free concentration of an analyte, and A and D are fluorochromes, the emission wavelength of fluorochrome D overlapping the excitation wavelength of fluorochrome A and the distance between A and D being in sufficiently close proximity to allow radiationless transfer of energy between the fluorochromes. A and D can be selected from a variety of acceptor, donor pairs such as fluorescein and tetramethylrhodamine.</p>		

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DETECTION OF ANALYTES
USING FLUORESCENT ENERGY TRANSFER

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5 Institute of General Medical Science and Grant No. NS27177 awarded by the National Institute of Neurological and Communicable Diseases. The Government has certain rights in this invention.

BACKGROUND OF THE INVENTION

10 This invention relates to energy transfer between fluorochrome labeled proteins to determine analyte concentration and, more particularly, to methods of determining cAMP concentration using labeled cAMP dependent protein kinase.

15 The function of many biological molecules is mediated through interactions with proteins and protein complexes. Many of these molecules are found in extremely small concentrations within a living cell. However, their concentration may transiently change in response to
20 physiological stimuli. In many instances, it is important to determine the concentration of such molecules to assess the metabolic state of a cell, hormonal responses and certain diseases. Even if the level changes several fold, at such minute concentrations, such change is not
25 detectable.

One such important molecule which mediates biological function through protein interaction is cyclic AMP (cAMP, adenosine 3', 5' - cyclic monophosphate). Cyclic AMP regulates intracellular reactions in all procaryotic and
30 nucleated animal cells that have been studied to date. It functions as a ubiquitous intracellular mediator, or second

messenger, of a variety of hormone induced effects, the first messenger being the extracellular hormone. Such hormone induced effects mediated by increased cAMP levels include, for example, triglyceride breakdown in a fat tissue induced by epinephrine, adrenocorticotrophic hormone (ACTH), glucagon or thyroid-stimulating hormone (TSH); water resorption in the kidney by vasopressin; glycogen breakdown in the muscle and liver; increased heart rate by epinephrine and progesterone secretion by the ovary in response to luteinizing hormone. These differential responses to cAMP are induced by hormone-receptor interactions at the cell surface and result in the intracellular synthesis of cAMP. Thus, different target cells respond to increases in cAMP levels in different but characteristic way.

For cAMP to function as a second messenger, its intracellular concentration must be tightly controlled and able to change rapidly in response to hormone-receptor binding. Normally, cAMP levels are less than about 1 μM and increase approximately five-fold upon hormonal stimulation. The increase in cAMP levels is due to synthesis by the enzyme adenylate cyclase from ATP. Following this transient increase, cAMP levels are rapidly returned to normal levels through the action of phosphodiesterases.

To effectively study cAMP dependent hormonal responses and their physiological mechanisms, it is necessary to accurately measure small concentrations of cAMP. More essentially, it is important to measure free cAMP concentrations, since in general, the biological activity of intracellular messengers is controlled by and correlates with the concentrations of free messenger, whereas total concentrations measured after tissue disruption include material bound to biologically irrelevant sites. Previous methods for measuring cAMP concentrations employed

classical competitive binding assays such as radioimmunoassays, and required the manipulation and preparation of extracts from thousands or millions of cells. These methods resulted in poor spatial and temporal
5 resolution and were incapable of distinguishing free or biologically active cAMP levels from total or sequestered cAMP. Moreover, these methods required the use of hazardous, perishable isotopes and were therefore expensive.

10 Additionally, second messengers other than cAMP such as cyclic GMP (cGMP), calcium and diacylglycerol as well as other organic molecules such as tumor-promoting phorbol esters are of great importance in signal transduction and cell physiology. Accurate determination of the
15 biologically active concentrations of these molecules is also necessary to effectively study hormonal responses and physiological mechanisms. However, except for simple inorganic ions such as calcium, there is no indicator system such as to measure the biologically active
20 concentrations of these molecules. Most of the same drawbacks that exist for determining free cAMP concentrations within the cell exist for these molecules as well.

Thus, there exists a need for a method to rapidly,
25 efficiently and non-destructively measure free intracellular cAMP concentrations as well as other organic molecules in single living cells. Such an invention would be of critical importance to the understanding of hormonal regulation. The present invention satisfies this need and
30 provides related advantages as well.

SUMMARY OF THE INVENTION

The invention provides labeled proteins suitable for determining the presence of cAMP, other second messengers

and organic molecules. The proteins are separately labeled with fluorochromes which, when in close spatial proximity, preferably, less than about 6 nm, interact through the transfer of energy from one fluorochrome to the other.

5 A composition of matter, $(S_1 \kappa A)_{n1}(S_2 \kappa D)_{n2}$ is provided wherein S_1 and S_2 are two proteins which are associated in one state and substantially disassociated in another, the equilibrium between which is controlled by the free
10 concentration of an analyte, and A and D are fluorochromes, the emission wavelength of fluorochrome D overlapping the excitation wavelength of fluorochrome A and the distance between A and D being in sufficiently close proximity to allow the radiationless transfer of energy between the fluorochromes. A and D can be selected from a variety of
15 acceptor, donor pairs such as fluorescein and tetramethylrhodamine and derivatives thereof. The concentration of analytes such as cAMP in a sample can be determined by contacting the sample with $(S_1 \kappa A)_{n1}(S_2 \kappa D)_{n2}$ providing energy near the excitation wavelength of D and
20 measuring the fluorescence of A or D, the concentration of cAMP and other said analytes being determined by the ratio of emission of D to the emission of A, previously calibrated with reference solutions of known analyte concentration.

25 BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 is a schematic diagram depicting the use of fluorescence energy transfer to measure cAMP.

Figure 2 shows the emission spectrum of labeled cAMPdPK at various concentrations of cAMP and the inset
30 plots the ratio of emission at 520 nm to 580 nm at the various cAMP concentrations.

Figure 3 shows the increase in cAMP concentration when a smooth muscle cell is treated with a membrane permeable cAMP analog.

Figure 4 shows the measurement of cAMP within a single cell when treated with various β_2 -adrenergic agonists and antagonists.

DETAILED DESCRIPTION OF THE INVENTION

As used herein, the term "protein" or "subunit" indicates a linear sequence of amino acids which exhibits biological function. This linear sequence includes native proteins, domains of proteins and fragments of proteins so long as function is maintained. The term also includes polypeptides and peptides.

As used herein, the term "associated" refers to two proteins such as subunits of an enzyme, which are in contact with each other. The contact region may include all or parts of the two molecules. Therefore, the term "substantially disassociated" or "disassociated" refers to the loss of contact between the associated regions, including loss of all contact regions such that the proteins are completely separated as well as the loss of some regions so that the body of the proteins are no longer in close proximity to each other but may still be tethered together. The term "complex" as used herein, refers to the proteins or subunits when they are in the associated state. The complex may include more than one molecule of each of the two proteins.

As used herein, the term "cAMPdPK" refers to the cyclic AMP dependent protein complex as described in Alberts et al., (1989), Molecular Biology of the Cell, Garland Publishing, Inc., and in Slice and Taylor, (1989),

J. Biol. Chem. 264:20940-20946, both of which are herein incorporated by reference. The complex consists of two proteins which perform regulatory and catalytic functions. It is intended that the term will include all isotypes of the regulatory and catalytic proteins.

As used herein, the term "fluorochrome" refers to a molecule which is capable of absorbing energy at a certain wavelength range and releasing the energy at a wavelength range other than the absorbance range. Therefore, the term "excitation wavelength" refers to the range of wavelengths that a fluorochrome absorbs energy. The term "emission wavelength" includes to the range of wavelengths that a fluorochrome releases energy or fluoresces. The term fluorochrome also includes derivatives thereof, such as chemical moieties for attachment to proteins. Examples of fluorochromes are fluorescein and tetramethylrhodamine. A derivative of each is fluorescein isothiocyanate and tetramethylrhodamine isothiocyanate.

As used herein, the term "sample" refers to a material, such as a solution, extract or intact cell suspected of containing an analyte.

As used herein, the term "labeling" refers to the attachment of a chemical moiety such as a fluorochrome to a protein. The attachment is typically through a covalent bond but may also include non-covalent interactions as well. Therefore, the term "labeled proteins" refers to proteins which have chemical moieties such as fluorochromes attached to them.

The invention provides a composition of matter, comprising $(S_1 \times A)_{n1}$ $(S_2 \times D)_{n2}$ wherein S_1 and S_2 are two proteins which are associated in one state and substantially disassociated in another state, and D and A are different fluorochromes, the emission wavelength of one fluorochrome

overlapping the excitation wavelength of the other fluorochrome, the distance between D and A is in sufficiently close proximity to allow the radiationless transfer of energy between the fluorochromes, and n1 and n2
5 representing the stoichiometry of said proteins, and wherein S_1 or S_2 is capable of binding an analyte X, the binding of X affects the affinity of S_1 and S_2 for each other.

The two proteins S_1 and S_2 which are labeled with
10 fluorochromes are typically subunits of an enzyme but may include any two proteins which exhibit associated and substantially disassociated states. Examples of such protein pairs include the subunits for cAMP dependent protein kinase (cAMPdPK), the subunits of GTP-binding
15 proteins, calmodulin and calmodulin dependent protein kinases and the regulatory and catalytic domains of protein kinase C.

When in the associated state, the subunits form a protein complex such as a holoenzyme. The number of
20 proteins within the complex can vary depending on the stoichiometry between S_1 and S_2 . Additionally, the complex may include proteins other than S_1 and S_2 . These other proteins within the complex may be considered as components of S_1 or S_2 so long as they maintain a stable association
25 with either S_1 or S_2 .

When in the disassociated state, at least one of the proteins will have substantially changed its distance from the other protein. For example, the disassociated state for cAMPdPK consists of free catalytic and regulatory
30 proteins or subunits. Likewise, GTP-binding proteins are disassociated into alpha and beta-gamma subunits, calmodulin dependent protein kinases disassociate from calmodulin, and protein kinase C regulatory and catalytic domains initially move apart and are subsequently separated

by proteolysis.

Radiationless energy transfer is based on the biophysical properties of fluorochromes. These principles are reviewed in Lakowicz, J., 1983, Principles of Fluorescence Spectroscopy, Plenum Press, New York and Jovin and Jovin, 1989, in Cell Structure and Function by Microspectrofluorometry, edited by E. Kohen and J. G. Hirschberg, Academic Press, both of which are incorporated herein by reference. Briefly, a fluorochrome absorbs light energy at a characteristic wavelength. This wavelength is also known as the excitation wavelength. The energy absorbed by a fluorochrome is subsequently released through various pathways, one being emission of photons to produce fluorescence. The wavelength of light being emitted is known as the emission wavelength and is an inherent characteristic of a particular fluorochrome. Radiationless energy transfer is the quantum-mechanical process by which the energy of the excited state of one fluorochrome is transferred without actual photon emission to a second fluorochrome. That energy may then be subsequently released at the emission wavelength of the second fluorochrome. The first fluorochrome is generally termed the donor (D) and has an excited state of higher energy than that of the second fluorochrome, termed the acceptor (A). The essential features of the process are that the emission spectrum of the donor overlap with the excitation spectrum of the acceptor, and that the donor and acceptor be sufficiently close. The distance over which radiationless energy transfer is effective depends on many factors including the fluorescence quantum efficiency of the donor, the extinction coefficient of the acceptor, the degree of overlap of their respective spectra, the refractive index of the medium, and the relative orientation of the transition moments of the two fluorochromes, but is typically 4-6 nm for favorable pairs of donor and acceptor. Beyond the optimum range of

intermolecular distances, the energy transfer efficiency falls off as the inverse sixth power of the distance.

The analyte X capable of being bound by either S_1 or S_2 , can be, for example, a second messenger such as cAMP, calcium, cGMP, and diacylglycerol; relevant hormone-receptor complexes; or an organic molecule such as phorbol ester. When bound to such molecules, S_1 and S_2 may be in either the associated or disassociated states.

The invention provides a composition of matter wherein S_1 and S_2 are proteins, or subunits of a complex, such as a holoenzyme. Such holoenzymes can be, for example, cAMPdPK, GTP-binding proteins, calmodulin dependent protein kinase and protein kinase C.

In one embodiment, the invention also provides a composition of matter wherein S_1 is the regulatory subunit of cAMPdPK and S_2 is the catalytic subunit of cAMPdPK and are labeled with different fluorochromes such that the emission wavelength of one overlaps the excitation wavelength of the other.

For example, in the absence of cAMP, the regulatory (R) and catalytic (C) subunits of cAMPdPK have a stoichiometry of R_2C_2 as shown in figure 1. The complex is inactive as a kinase, since the R subunits inhibit the kinase domains of the C subunits. Upon the binding of cAMP by R, the R subunits dissociate from the C subunits, releasing the biological activity of the latter. The invention includes labeling the R subunits with one fluorochrome and the C subunits with a different fluorochrome, such that in the intact R_2C_2 complex, light energy absorbed by one fluorochrome is transferred to the other fluorochrome by the process of radiationless energy transfer. This phenomenon requires that the wavelengths of fluorescent emission from one fluorochrome, the donor (D),

overlap the wavelengths of excitation of the other fluorochrome, the acceptor (A), and that the D and A moieties be in sufficiently close proximity to allow the radiationless transfer of energy between the fluorochrome pairs. In the case of fluorescein and tetramethylrhodamine, this distance is less than about 4 to 6 nm. For other fluorochrome pairs, the distance can be determined by those skilled in the art. Provided suitable choices are made for D and A, this condition of close proximity is reasonably well satisfied in a holoenzyme complex such as that for CAMPdPK, so that energy transfer occurs with considerable efficiency.

The transferred energy is easily detectable by excitation at wavelengths absorbed by D, since emission from D is quenched and the energy appears instead at longer wavelengths characteristic of A's emission.

Subunits RkD and CkA can be paired to make $(RkD)_2(CkA)_2$, or RkA and CkD can be combined into $(RkA)_2(CkD)_2$, where k denotes attachment of the indicated fluorochrome. Once CAMP has dissociated R from C, D and A thereby become widely separated from each other and unable to transfer energy from D to A. With the subunits dissociated, illumination at wavelengths absorbed by D simply causes fluorescence emission from D, with no involvement of A, so that the composite emission spectrum shifts towards the shorter wavelengths of D.

Holoenzyme labeled with two different fluorochromes as described above is found to have essentially identical sensitivity to CAMP as the native holoenzyme. This property allows the use of protein complexes such as CAMPdPK to be used as measuring devices or sensors for a variety of biological molecules. Moreover, it guarantees that the sensor has the optimal affinity for such biological molecules as CAMP, for example, since native

holoenzyme is the main endogenous sensor for cAMP and the concentrations of interest are precisely those which affect this enzyme. Moreover, the activity of labeled enzyme as a kinase upon stimulation by cAMP is found to be identical to native holoenzyme likewise stimulated. This property is valuable because it minimizes any perturbation of the biological effectiveness of cAMP within cells into which the sensor has been introduced. If the sensor were not an effective kinase, then any cAMP bound to it would be prevented from exerting biological activity so that the sensor would inherently inhibit the signaling pathway whose activity was to be measured. However, if the sensor is a normal kinase, cAMP molecules bound to it are biologically fully effective and not removed from mediating such events.

The invention also provides a method of determining the concentration of an analyte in a sample comprising the steps of: (a) labeling proteins S_1 and S_2 with fluorochromes A and D, where S_1 or S_2 is capable of binding the analyte and S_1 and S_2 are associated in one state and substantially disassociated in another state, the equilibrium of which is controlled by the concentration of free analyte, the fluorochromes are selected such that the emission wavelength of one fluorochrome overlaps the excitation wavelength of the other fluorochrome to form labeled proteins $(S_1D)_{n1}$ $(S_2A)_{n2}$ or $(S_1A)_{n1}$ $(S_2D)_{n2}$; (b) contacting the labeled proteins with the sample; (c) providing energy near the excitation wavelength of D or A; (d) measuring the fluorescences emitted at the emission wavelengths of A and D respectively, and determining the extent of energy transfer from D to A by the ratio of their emission amplitudes, or by the fluorescence lifetime of D, or by the rate of photobleaching of D. Any of these parameters, but most conveniently the ratio of emission amplitudes, can be related to the analyte concentration by a calibration curve established with samples of known analyte concentration.

In one embodiment, holoenzyme labeled with fluorescein on the C subunit and tetramethylrhodamine on the R subunit, for example, can be used to measure the concentration of cAMP in vitro and in vivo. For the former, the enzyme is

5 titrated with cAMP while measuring its fluorescent emission following excitation at 490 to 500 nm. A plot of the emission ratio (520:580 nm are the optimal wavelengths) against known concentrations of cAMP can be used as a standard curve for determining cAMP concentrations in

10 unknown solutions under the same conditions. To measure in vivo cAMP concentrations inside living cells requires incorporation of the labeled holoenzyme into the cells of interest by microinjection, electroporation, lipofection, bead loading, scrape loading or any other comparable and

15 well-known technique. Such methods are described in Riabowol et al., (1988) Cold Spring Harbor Symposia on Quantitative Biology 53:85-90; Sambrook et al., Molecular Cloning: A Laboratory Manual, (1989) Cold Spring Harbor Laboratory and Ausubel et al., Current Protocols in

20 Molecular Biology, (1987), John Wiley and Sons, all of which are incorporated herein by reference. Excitation at 490 to 500 nm on a fluorescence microscope and detection of the emission near the two optimal wavelengths (520 and 580 nm) is achieved by a photometer or television camera.

25 Calibration of the minimum and maximum ratio value is achieved by first setting the zero cAMP concentration value by addition of a cAMP antagonist, adenosine -3', 5'- cyclic monophosphothioate, Rp-isomer (Rp-cAMPS) which inhibits dissociation of the holoenzyme. Subsequent removal of the

30 cAMP antagonist and addition of the membrane-permeable cAMP analog gives complete dissociation and hence intermediate cAMP values can be determined using the standard curve derived in vitro.

Currently preferred choices of fluorochromes are

35 fluorescein for the donor dye, which has emission wavelengths of about 510 to 550 nm, and

tetramethylrhodamine for the acceptor, which has excitation wav lengths of about 510 to 550 nm, both being attached to the subunits via isothiocyanates under conditions to favor reaction with lysines. However, a variety of fluorochromes and derivatives thereof can be used so long as the emission wavelength of fluorochrome D overlaps the excitation wavelength of fluorochrome A and results in energy transfer from D to A. For example, D and A fluorochrome pairs may be a 7 aminocoumarin and dipyrromethene-borate, an indocarbocyanine and rhodamine X or a phycoerythrin and allophycocyanin.

The invention also provides for a method of preparing a complex comprising protein subunits S_1 and S_2 which are associated in one state and substantially disassociated in another state, the equilibrium between which is controlled by the concentration of a free analyte X, and wherein S_1 or S_2 is capable of binding the analyte, comprising: (a) attaching fluorochrome A to S_1 , said fluorochrome A having an excitation wavelength which overlaps with the emission wavelength of a second fluorochrome D; (b) attaching the second fluorochrome D to S_2 ; (c) mixing the labeled subunits so that a complex is formed wherein the fluorochromes are in sufficiently close proximity to allow radiationless transfer of energy between the fluorochromes; and (d) recovering the labeled subunits as a complex.

In one embodiment, R and C are separately available as recombinant proteins, for example from expression in E. coli, which are then labeled. However, careful adjustment of reaction conditions is necessary so that the act of labeling does not prevent reassociation into holoenzyme or subsequent cAMP sensitivity and kinase activity. Previous literature has suggested that labeling isolated C with fluorescein isothiocyanate ruined its ability to combine with R; under milder conditions, however, C was not damaged as described below.

The invention additionally provides a method of preparing a complex comprising protein subunits S_1 and S_2 which are associated in one state and substantially disassociated in another state, the equilibrium between which is controlled by the concentration of a free analyte X, and wherein S_1 or S_2 is capable of binding the analyte, comprising: (a) attaching fluorochrome A to a complex comprising S_1 and S_2 , said fluorochrome A having an excitation wavelength which overlaps with the emission wavelength of a second fluorochrome D; (b) attaching the second fluorochrome D to a second complex comprising S_1 and S_2 ; (c) separating the subunits of said first and second complex to obtain labeled subunits S_{1KD} , S_{2KD} , S_{1KA} and S_{2KA} ; and (d) mixing the labeled subunits wherein said fluorochromes are in sufficiently close proximity to allow radiationless transfer of energy between the fluorochromes to form label complexes $(S_{1KD})_{n1}$ $(S_{2KA})_{n2}$ or $(S_{1KA})_{n1}$ $(S_{2KD})_{n2}$, wherein $n1$ and $n2$ represent the stoichiometry of the labeled subunits.

An alternative method is to label pre-formed holoenzyme R_2C_2 with D, thereby preparing $(R_{KD})_2(C_{KD})_2$ in which the sites of interaction between R and C are inherently protected. A separate batch of R_2C_2 is analogously labeled with A to make $(R_{KA})_2(C_{KA})_2$. Then labeled R and labeled C are dissociated and separated from each other by passage through a cAMP affinity column, to which labeled R sticks while labeled C passes through. R can be recovered from the column by elution with free cAMP. When this separation is applied to each batch, one obtains R_{KD} , C_{KD} , R_{KA} , and C_{KA} , which are then recombined in different pairings to make $(R_{KD})_2(C_{KA})_2$ as one preparation and $(R_{KA})_2(C_{KD})_2$ as another.

The holoenzyme or individual subunits of cAMPdPK can be prepared from mammalian sources such as that described by Bubis et al., (1985) Biochemistry, 24:2163-2170, and

Zick et al., J. Biol. Chem., (1982) 257:2287-2293, both of which are herein incorporated by reference. If subunits from recombinant or mammalian sources are used for the preparation of holoenzyme then reconstitution can be performed as described in the Examples.

The following examples are intended to illustrate but not limit the invention.

EXAMPLE I

Recombinant Expression and Purification 10 of cAMP-Dependent Protein Kinase Regulator Subunit

Reagents were purchased from the following companies: histone IIA, phenylmethylsulfonyl fluoride, isopropyl- β -D-thiogalactopyranoside (IPTG), and bovine serum albumin, (Sigma, St. Louis, MO); N^{α} -p-tosyl-L-lysine chloromethyl ketone and cAMP, (United States Biochemical Corp., 15 Cleveland, OH); 2,8- $[^3\text{H}]$ cAMP (27 Ci/mmol) and 8- N_3 - $[^{32}\text{P}]$ cAMP (20-30 Ci/mmol), (ICN); Sephadex G-150 and cAMP-agarose (N^6 -ethane spacer), (Pharmacia, Pleasant Hill, CA); nitrocellulose (0.45 μm) and electrophoresis reagents, 20 (Bio-Rad, Richmond, CA); and Cytoscint, (West Chem). Enzymes used in DNA manipulations were obtained from either Bethesda Research Laboratories (Gaithersburg, MD) or Boehringer Mannheim (Indianapolis, IN) and were used according to the manufacturer's specifications. 25 Fluorescent labels were obtained from Molecular Probes (Eugene, OR) and cAMP analogs were obtained from Biolog (La Jolla, CA). All other reagents were analytical grade.

Plasmid 62C12, Lee et al., (1983) Proc. Natl. Acad. Sci. U.S.A., 80:3608-3612, and herein incorporated by 30 reference, which contains the full-length cDNA of bovine R^1 in pBR322 was subcloned into pUC7, Viera et al., (1982) Gene, 19:259-268, and herein incorporated by reference for cDNA expression.

For subcloning, DNA manipulations were carried out according to Maniatis et al., (1982) Molecular Cloning: A Laboratory Manual, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, and herein incorporated by reference.

5 Briefly, 62C12 was digested with NarI, and the 3.6-kilobase (kb) fragment containing the cDNA was isolated. Following partial digestion with NcoI, the two large fragments, the full-length cDNA fragment (1155) base pairs (bp), and a truncated fragment (1020 base pairs) were isolated as a

10 mixture, filled-in with Klenow fragment, and ligated into pUC7 that had been linearized with Hinc II. The 135-base pair fragment also was isolated, radiolabeled with [γ - 32 P]ATP, and used to identify those clones which contained the full-length insert. Twenty colonies were selected, and

15 the DNA from each was prepared and digested with Bg/I in order to establish the orientation of the insert. The DNA from three clones gave a restriction pattern which indicated that the R' coding segment was inserted in the correct orientation with regard to the lac Z' gene. These

20 clones were further characterized for expression of the protein.

Since basal levels of cAMP are high in E. coli, the following method was devised for the accurate detection of R in crude extracts independent of exogenous cAMP. This

25 method could be used for rapid screening. Protein samples were electrophoresed on 12.5% polyacrylamide gels (1.5 mm) containing sodium dodecyl sulfate according to the method of Laemmli (1970) Nature 227:680-685. Proteins were then transferred from the gels to nitrocellulose filters using

30 an electroblotting apparatus (Hoefer Scientific Instruments). Gels were electrotransferred for 4 to 6 hours at 500 mA in 20 mM Tris, 154 mM glycine, 20% methanol (pH 8.3). The nitrocellulose was incubated for 1 hour at room temperature with 0.05% Tween 20 in 150 mM sodium

35 chloride and 10 mM Tris (pH 7.5) and then washed at room temperature with the same buffer except that Tween 20 was

omitted. The filter was incubated with 8-N₃[³²P]cAMP (20 nM) in the same buffer for 30 minutes in the dark at room temperature. After washing with ice-cold buffer, the filter was irradiated with UV S-11 lamp (254 nm) for 5 minutes, washed again, blotted dry, and subjected to autoradiography. The R-subunit also was detected by Millipore filtration using 2,8-[³H]cAMP as described by Kerlavage et al., (1982) J. Biol. Chem., 257:1749-1754, and herein incorporated by reference.

10 Purification of the regulatory subunit was performed by inoculating 5 mls of L-broth containing ampicillin (50 µg/ml) with the highest expressing transformant and growing at 37°C until the cell density reached OD₅₅₀ = 0.1-0.5. The culture was then transferred to 1 liter of L-broth
15 containing ampicillin and incubated (12 to 16 hours). Unless indicated, all subsequent procedures were carried out at 4°C. The cells were centrifuged at 5,000 x g for 30 minutes, and the resulting pellet was resuspended in buffer I (20 mM potassium phosphate, 5 mM EDTA, 5 mM 2-
20 mercaptoethanol, 15 mg/liter N^α-p-tosyl-L-lysine chloromethyl ketone, and 15 mg/liter phenylmethylsulfonyl fluoride). This suspension was passed through a French pressure cell twice and then centrifuged at 5,000 x g for 20 minutes. The pellet was re-extracted with the same
25 buffer. Ammonium sulfate was added slowly to the pooled supernatant fractions until a concentration of 70% was reached. After 1 hour, the precipitated protein was recovered by centrifugation at 12,000 x g for 30 minutes, redissolved in buffer I, and dialyzed against the same
30 buffer overnight. Fractions were assayed for [³H]cAMP binding and analyzed by polyacrylamide gel electrophoresis followed by photolabeling as described above. The dialyzed sample was then added to cAMP-agarose and gently rotated overnight. After removing the supernatant solution, the
35 resin was washed with buffer I containing 2 M NaCl until the absorbance at 280 nm reached 0 and then washed again

with buffer I. The R-subunit was eluted with 2 volumes of CAMP (50 mM) in buffer I at 30°C for 1 hour. The elution was repeated at least once.

The eluate from the CAMP-agarose resin frequently
5 contained some proteolytic degradation products of R in addition to full-length R, which was separated on a 70 cm Sephadex G-150 column equilibrated and eluted with 25 mM potassium phosphate, 5 mM 2-mercaptoethanol, 5 mM EDTA, and 150 mM NaCl (pH 6.5). Fractions (2 ml) were collected at
10 a flow rate of 7 ml/hour. The R-subunit eluted close to the void volume and was well-resolved from the 35-kDa proteolytic fragment.

The recombinant CAMP dependent regulatory subunit was sequenced on an Applied Biosystems Gas-phase Sequencer.
15 Two nanomoles of R-subunit were sequenced, and the phenylthiohydantoin derivatives were analyzed on a Beckman HPLC system using a cyano column (IBM) according to the procedure of Hunkapillar and Hood, (1983) Methods Enzymol., 91:486-493, and herein incorporated by reference.

20

EXAMPLE II

Recombinant Expression and Purification of
CAMP-Dependent Protein Kinase Catalytic Subunit

Plasmid containing the full length clone for the α catalytic subunit of murine CAMP-dependent protein kinase,
25 pMCO, Uhlen et al., (1986) Proc. Natl. Acad. Sci. USA, 83:1300-1304, which is incorporated herein by reference, was subcloned into M13 mp18 for mutagenesis. The bacterial expression vector, pT7-7, Tabor and Richardson, (1985) Proc. Natl. Acad. Sci. USA, 82:1074-1078, and herein
30 incorporated by reference was used for recombinant expression. The E. coli strains used to express the catalytic subunit were: JM101, 222, K38 and BL21 (DE3).

B fore introducing a unique NdeI restriction site into the catalytic subunit gene at the initiation methionine, the 1.9-kilobase SacI fragment from pMCO containing the coding region for the catalytic subunit was excised and
5 religated into the SacI site of M13mp18. Single stranded DNA was prepared and used as template for oligonucleotide-directed mutagenesis with a 24-base oligonucleotide (5'-CACGCCGCCCATATGGGCAACGCC-3') according to the procedure described by Zoller and Smith, (1984) DNA 3:479-488, and
10 herein incorporated by reference. The mutation was verified by dideoxy sequencing according to Sanger et al., (1980) J. Mol. Biol. 143:161-178, and herein incorporated by reference and by mapping with restriction enzymes.

Construction of pLWS-3 was performed by excising a
15 1988-base fragment from the M13 construct described above with NdeI and HindIII, purified from an agar gel, and ligated into the NdeI-HindIII restriction sites of pT7-7. This ligation mixture was used to transform E. coli JM101 using the procedure described by Maniatis et al., *ibid*.

20 Transformants were selected and screened for plasmid DNA containing the cDNA insert following digestion with NdeI and HindIII. DNA was isolated from this plasmid, pLWS-3, and the sequence confirmed by mapping with restriction enzymes and by sequence analysis. The plasmid
25 was then used to transform E. coli BL21(DE3), and transformants were screened for expression of the catalytic subunit by monitoring for the appearance of a 38-40 kDa band following polyacrylamide gel electrophoresis.

For expression of the catalytic subunit, one liter of
30 L-broth (10% Tryptone, 5% yeast extract, and 5% NaCl) containing ampicillin was inoculated with 5 ml of a stationary phase culture of BL21(DE3) transformed with pLWS-3 and grown at 37°C with vigorous shaking until an absorbance of 0.5 at 590 nm was reached. After adding

isopropyl- β -D-thiogalactopyranoside to a final concentration of 0.5 mM, the cells were grown for an additional 3 hours at 37°C and then harvested by centrifugation.

5 Purification of the recombinant catalytic subunit was performed by resuspending the pellet in 6 ml of lysis buffer (30 mM MES, pH 6.45, 50 mM KCl, 1 mM EDTA, and 5 mM β -mercaptoethanol) at 4°C and passed twice through a French pressure cell. Additional lysis buffer was added to bring
10 the total volume to 30 ml. After centrifugation at 12,000 x g for 15 minutes at 4°C, the supernatant was removed and diluted with ice-cold water until the conductivity was below 1.3 mmhos. The diluted supernatant was applied to a phosphocellulose column (Whatman P11, 2.0 x 8.0 cm) pre-equilibrated with buffer I (30 mM MES, pH 6.45, 1 mM EDTA).
15 After washing with the equilibration buffer until the absorbance at 280 nm was 0, the catalytic subunit was eluted with a linear gradient of buffer I (50 ml) to 0.5 M potassium phosphate, pH 7.0 (50 ml). Two-ml fractions were
20 collected, and those containing kinase activity were pooled. The protein was precipitated by the batchwise addition of solid NH_4SO_4 , with constant stirring, until the solution was 80% saturated. The precipitate was collected by centrifugation at 12,000 x g for 20 minutes and then
25 resuspended in 0.8 ml of H_2O . After removing the undissolved material by centrifugation, the protein solution was loaded onto an Ultrogel ACA 44 (LKB) gel filtration column (1.0 x 116 cm) equilibrated with 25 mM potassium phosphate, pH 6.8. The fractions containing
30 kinase activity were combined and concentrated to 1 mg/ml by ultrafiltration using an Amicon apparatus and a PM-30 membrane. Based on polyacrylamide gel electrophoresis carried out in the presence of sodium dodecyl sulfate (SDS) according to Laemmli, (1970) Nature 227:680-685, which is
35 incorporated herein by reference, the recombinant catalytic subunit was nearly homogeneous, and the yield was 3.5 to

4.0 mg/liter of culture.

Recombinant catalytic subunit was diluted to 0.04 mg/ml with 20 mM Tris, pH 7.5. Fifty- μ l aliquots were kept on ice and then placed in a 49°C water bath. At the
5 indicated times, tubes were removed, placed on ice, and assayed for stability of the recombinant protein.

Isoelectric focusing was carried out using a LKB multiphor model 2117 with pre-cast LKB Ampholine Pagplates, pH. 3.5 to 9.5. The standards were bovine carbonic
10 anhydrase (pI 6.57), myoglobin (pI 6.67 and 7.16), and lactate dehydrogenase (pI 8.55).

The molecular weight of the recombinant catalytic subunit was determined by gel filtration using a LKB Ultrogel AcA 44 column (1 x 116 cm). The running buffer
15 was 25 mM potassium phosphate (pH 6.8), 200 mM KCl, and 0.2% sodium azide. The standards and their respective M values were ribonuclease A (13,700), chymotrypsinogen A (25,000), porcine catalytic subunit (38,000), ovalbumin (43,000), and bovine serum albumin (67,000).

20 The recombinant catalytic subunit was sequenced on a Applied Biosystems gas-phase sequencer as described in Example I.

EXAMPLE III

Purification of Mammalian Catalytic Subunit

25 Porcine hearts (40 hearts) were homogenized and then centrifuged. The supernatant (conductivity 3 mmho) was passed directly through a DE-23 column (4 L resin) equilibrated in 15 mM potassium phosphate, pH 6.5, 2 mM
EDTA, 5 mM β -mercaptoethanol. The protein was eluted from
30 the resin with buffer E (40 mM potassium phosphate, pH 6.5, 2 mM EDTA, 5 mM β -mercaptoethanol). The pooled kinase peak

was immediately adjusted to a conductivity of 4 mmho at which point 1.5 L of DE-52 resin was added and the whole was mechanically stirred for 2 hours. The supernatant was poured off and the resin was poured into a column and washed extensively with buffer E, followed by extensive washing with buffer F (17 mM potassium phosphate, pH 6.1, 2 mM EDTA, 5 mM β -mercaptoethanol). The C-subunit was then eluted with 3 L of 220 μ M CAMP followed by 8 L of buffer F and the eluent was passed through a CM-Sepharose column (50 mL volume) and then eluted with a linear gradient of buffer F containing 10% glycerol (250 mL) to 200 mM potassium phosphate, pH 6.1, 10% glycerol, 2 mM EDTA, 5 mM β -mercaptoethanol (250 mL).

EXAMPLE IV

Preparation of Labeled cAMP-Dependent Protein Kinase

(a) Labeling subunits:

Recombinant catalytic and regulatory subunits (approximately 0.5-2.0 mg/ml) of mammalian cAMP-dependent protein kinase were purified as described in Examples I and II and were dialyzed separately against 25 mM bicine pH 8.0, 0.1 mM EDTA for 4 to 5 hours at 0°C. The catalytic subunit was labeled with 0.3 mM fluorescein 5'-isothiocyanate (FITC; Molecular Probes, Eugene, OR) in the presence of 8 mM $MgCl_2$ and 5 mM ATP to prevent inactivation of kinase activity. The regulatory subunit was labeled with 0.5 mM tetramethylrhodamine isothiocyanate (TRITC, isomer G; Molecular Probes, Eugene, OR). Both labelings were allowed to proceed from 30 minutes at room temperature, then quenched by addition of 5 mM glycine for 10 to 15 minutes. The excess dye was removed by passing each protein solution through a Sephadex G-25 column (3 ml), eluted with 25 mM potassium phosphate pH 6.8, 2 mM EDTA, 5 mM 2-mercaptoethanol, 10% glycerol. The first colored or fluorescent band was collected.

The FITC-labeled catalytic and TRITC-labeled regulatory subunits (equal amounts by weight) were then mixed and dialyzed against 25 mM potassium phosphate pH 6.8, 0.5 mM $MgCl_2$, 0.1 mM ATP, 5 mM 2-mercaptoethanol, 5% glycerol for 3 to 5 days at 4°C. The formation of holoenzyme could be demonstrated by kinase assay measured using Kemptide as substrate (Slice & Taylor, *ibid.*).

(b) Labeling holoenzyme:

cAMP-dependent protein kinase holoenzyme (0.5 - 1.0 mg/ml) prepared from mammalian catalytic and recombinant regulatory subunits was dialyzed against 25 mM bicine pH 8.0, 0.1 mM EDTA at 4°C. $MgCl_2$ and ATP were added to give final concentrations of 8 mM and 5 mM respectively. The proteins were labeled with either fluorescein 5'-isothiocyanate or tetramethylrhodamine isothiocyanate (Isomer G) at final concentrations of 0.1 - 0.5 mM at room temperature for 20 to 30 minutes, followed by quenching with 5 mM glycine pH 8.0 for 10 minutes. Excess free dye was (partially) removed by passing the protein solutions through a column of Sephadex G-25 (3 ml), eluting with 10 mM potassium phosphate pH 6.7, 150 mM KCl, 5 mM 2-mercaptoethanol, 0.1 mM EDTA (Buffer I) and collecting the first colored fluorescent band.

The labeled holoenzymes were then passed through a cAMP affinity column (0.25 ml) (Wolfgang, W. et al., *FEBS Lett.* 99, 62 (1979)) at a rate of approximately 0.1 ml/minute and then eluted with 2 column volumes of the Buffer I at 4°C. The collected fractions contained labeled catalytic subunit. The column was eluted with Buffer I containing 1 M KCl (2 column volumes) and the Buffer I (2 column volumes) again. These washes may contain some additional labeled catalytic subunit. To recover the labeled regulatory subunit, the column was eluted slowly with 30 mM cAMP in Buffer I (4 column volumes) at room

temperature over a 2 hour period.

The differently labeled subunits were recombined in appropriate donor-acceptor pairs and dialyzed against several changes of 25 mM potassium phosphate pH 6.8, 0.5 mM MgCl₂, 0.1 mM ATP, 5 mM 2-mercaptoethanol, 5% glycerol for 3 to 5 days at 4°C. The formation of cAMPdPK holoenzyme could be monitored by loss of kinase activity, which can be recovered on addition of cyclic AMP.

EXAMPLE V

10 Titration of cAMP With Labeled Holoenzyme In Vitro

Holoenzyme (20 nM) labeled with 5'-fluorescein isothiocyanate (5'-FITC) on the catalytic subunit and tetramethylrhodamine isothiocyanate (isomer G) on the regulatory subunit, was dissolved in 135 mM KCl, 10 mM K₂MOPS pH 7.2, 2 mM MgCl₂, 1 mM ATP at room temperature. The fluorescence emission was measured following excitation at 495 nm before and after addition of 20, 50, 100, 200, 400 nM, 1 and 51 μM cAMP (figure 2). Only the 0, 50, 200, 1000 nM cAMP spectra are shown in Figure 1 for clarity. The calculated ratio of emission at 520 nm to 580 nm, normalized to the ratio at 0 cAMP is plotted with cAMP concentration in Figure 2 (inset), indicated a half-maximal change in ratio as about 100 nM cAMP.

EXAMPLE VI

25 In Vivo Measurement of cAMP in Single Smooth Muscle Cells

Smooth muscle cells (BC₃H₁ cell line) were microinjected as described by Riabowol, Gilman & Feramisco, Cold Spring Harbor Symposia on Quantitative Biology, 53:85-90 (1988), which is incorporated herein by reference, with 60 μM holoenzyme that was labeled with 5'-FITC on the C subunit and TRITC on the R^I subunit (figure 3). After 15 to

20 minutes, the cells were illuminated on a fluorescence microscope stage with 490 nm light at room temperature. The emitted light was detected by a Silicone Intensified Target camera (Dage-NTI) after passing through either a 500-530 nm bandpass filter or a 580 nm longpass filter. The two pictures per time point were digitised by an image processor, the ratio of 500-530 nm to > 580 nm calculated at each pixel and displayed in pseudocolors on a television monitor. The average ratio in a representative cell was determined at each time point to give Figure 3. At time point 1, 250 μ M dibutyryl cAMP, a membrane-permeable cAMP analog, was added resulting in a gradual increase in ratio indicating a rise in cAMP analog inside the cell.

In a separate experiment BC_3H_1 cells were microinjected and the emission ratio determined as described above. The following additions were made and are shown in figure 4 (1) 100 nM isoproterenol (2) 100 nM propranolol (3) 50 μ M forskolin (4) 250 μ M dibutyryl cAMP. Addition of isoproterenol, a β_2 -adrenergic agonist elicited a rapid rise in emission ratio and therefore cAMP concentration within the cell, which was subsequently decreased by propranolol (a β_2 -adrenergic antagonist). Direct activation of adenylate cyclase by forskolin caused a maximal emission ratio as no further effect was seen with dibutyryl cAMP, a membrane-permeable cAMP analog.

Although the invention has been described with reference to the presently-preferred embodiment, it should be understood that various modifications can be made without departing from the spirit of the invention. Accordingly, the invention is limited only by the following claims.

WE CLAIM:

1. A composition of matter, comprising $(S_1 \times A)_{n1}$
 $(S_2 \times D)_{n2}$ wherein S_1 and S_2 are two proteins which are
associated in one state, and substantially disassociated in
another state the equilibrium between which is controlled
5 by the concentration of a free analyte X, and D and A are
different fluorochromes, the emission wavelength of one
fluorochrome overlapping the excitation wavelength of the
other fluorochrome, the distance between D and A being in
sufficiently close proximity to allow radiationless
10 transfer of energy between the fluorochromes, and $n1$ and $n2$
representing the stoichiometry of said proteins, and
wherein S_1 or S_2 is capable of binding the analyte.
2. The composition of matter of claim 1, wherein
 S_1 and S_2 are subunits of an enzyme.
3. The composition of matter of claim 2, wherein
the subunits are selected from the group of protein pairs
consisting of the subunits for cAMPdPK, the subunits of
GTP-binding protein subunits, calmodulin and calmodulin
5 dependent protein kinases and the regulatory and catalytic
domains of protein kinase C.
4. The composition of matter of claim 1, wherein
 S_1 is the regulatory subunit of cAMPdPK and S_2 is the
catalytic subunit of cAMPdPK.
5. The composition of matter of claim 1, wherein
D and A are selected from the group consisting of
fluorescein and tetramethylrhodamine, a 7 aminocoumarin and
dipyrrromethene-borate, an indocarbocyanine and rhodamine X
and a phycoerythrin and allophycocyanin, or derivatives
thereof, provided that the selected pair exhibit
fluorescence energy transfer.

6. The composition of matter of claim 1, wherein the emission wavelength of D and the excitation wavelength of A are about 510 to 550 nm.

7. The composition of matter of claim 5 or 6, wherein the fluorochromes are fluorescein and tetramethylrhodamine or derivatives thereof.

8. The composition of matter of claim 4, wherein n_1 is 2 and n_2 is 2.

9. The composition of matter of claim 1, wherein the molecule capable of being bound by S_1 or S_2 is selected from the group consisting of cAMP, GTP, hormone receptor complexes, calcium, diacylglycerol and phorbol esters.

10. The composition of matter of claim 1, wherein the analyte is cAMP.

11. The composition of matter of claim 1, wherein S_1 and S_2 are recombinant proteins.

12. The composition of matter of claim 1, wherein one protein is a recombinant protein and one protein is derived from mammalian sources.

13. The composition of matter of claim 1, wherein S_1 and S_2 are derived from mammalian sources.

14. A method of determining the concentration of an analyte in a sample comprising the steps of:

- 5 (a) labeling proteins S_1 and S_2 with fluorochromes A and D, where S_1 or S_2 is capable of binding the analyte and S_1 and S_2 are associated in one state and substantially disassociated in another state, the equilibrium of which is controlled by the concentration of free analyte, the fluorochromes characterized in that the emission wavelength of fluorochrome D overlaps the excitation wavelength of fluorochrome A to form labeled proteins
- 10 $(S_1KD)_{n1}$ $(S_1KA)_{n2}$;
- 15 (b) contacting the labeled proteins with the sample;
- (c) providing energy near the excitation wavelength of D or A; and
- 20 (d) measuring the fluorescence emitted at the emission wavelength of A, the extent of energy transfer from D to A being determined by the emission amplitude.

25 15. The method of claim 14, wherein the measuring step further comprises measuring the fluorescence emitted at the emission wavelength of the extent of energy transfer from D to A being determined by the ratio of the emission amplitudes of A and D.

30 16. The method of claim 14, wherein the measuring further step comprises measuring the fluorescence lifetime of D or the rate of photobleaching of D.

17. The method of claim 14, wherein the sample comprises an intact cell.

18. The method of claim 17, wherein the contacting step further comprises a method selected from the group consisting of microinjection, electroporation, lipofection, bead loading and scrape loading.

5 19. The method of claim 14, wherein the proteins S_1 and S_2 comprise subunits of an enzyme.

20. The method of claim 19, wherein the subunits are selected from the group of protein pairs consisting of the subunits for cAMPdPk, the subunits of GTP-binding
10 protein subunits, calmodulin and calmodulin dependent protein kinases and the regulatory and catalytic domains of protein kinase C.

21. The method of claim 14, wherein S_1 is the regulatory subunit of cAMPdPK and S_2 is the catalytic
15 subunit of cAMPdPK.

22. The method of claim 14, wherein D and A are selected from the group consisting of fluorescein and tetramethylrhodamine, a 7 aminocoumarin and dipyrrometheneborate, an indocarbocyanine and rhodamine X and a
20 phycoerythrin and allophycocyanin, or derivatives thereof, provided that the selected pair exhibit fluorescence energy transfer.

23. The method of claim 14, wherein the emission wavelength of D and the excitation wavelength of A are
25 about 510 to 550 nm.

24. The method of claim 22 or 23, wherein the fluorochromes are fluorescein and tetramethylrhodamine or derivatives thereof.

25. The method of claim 21, wherein n_1 is 2 and n_2 is 2.

26. The method of claim 14, wherein the analyte is selected from the group consisting of cAMP, GTP, hormone receptor complexes, calcium, diacylglycerol and phorbol esters.

27. The method of claim 14, wherein the analyte is cAMP.

28. The method of claim 14, wherein S_1 and S_2 are recombinant proteins.

29. The method of claim 14, wherein one protein is a recombinant protein and one protein is derived from mammalian sources.

30. The method of claim 14, wherein S_1 and S_2 are derived from mammalian sources.

31. A method of preparing a complex comprising protein subunits S_1 and S_2 which are associated in one state and substantially disassociated in another state, the equilibrium between which is controlled by the concentration of a free analyte X, and wherein S_1 or S_2 is capable of binding the analyte, comprising:

- 10 (a) attaching fluorochrome A to S_1 , said fluorochrome A having an excitation wavelength which overlaps with the emission wavelength of a second fluorochrome D;
- (b) attaching the second fluorochrome D to S_2 ;
- 15 (c) mixing the labeled subunits so that a complex is formed wherein said fluorochromes are in sufficiently close proximity to allow radiationless transfer of energy between the fluorochromes; and
- 20 (d) recovering the labeled subunits as a complex.

32. The method of claim 31, wherein S_1 is the regulatory subunit of cAMPdPK and S_2 is the catalytic subunit of cAMPdPK.

33. The method of claim 31, wherein the fluorochromes are fluorescein tetramethylrhodamine or derivatives thereof.

34. The method of claim 31, wherein the analyte is cAMP.

35. The method of claim 31, wherein S_1 and S_2 are recombinant proteins.

36. The method of claim 31, wherein one protein is a recombinant protein and one protein is derived from mammalian sources.

37. The method of claim 31, wherein S_1 and S_2 are derived from mammalian sources.

38. A method of preparing a complex comprising protein subunits S_1 and S_2 which are associated in one state and substantially disassociated in another state, the equilibrium between which is controlled by the concentration of a free analyte X, and wherein S_1 or S_2 is capable of binding the analyte, comprising:

- (a) attaching fluorochrome A to a complex comprising S_1 and S_2 , said fluorochrome A having an excitation wavelength which overlaps with the emission wavelength of a second fluorochrome D;
- (b) attaching the second fluorochrome D to a second complex comprising S_1 and S_2 ;
- (c) separating the subunits of said first and second complex to obtain labeled subunits $S_1\kappa D$, $S_2\kappa D$, $S_1\kappa A$ and $S_2\kappa A$; and
- (d) mixing the labeled subunits wherein said fluorochromes are in sufficiently close proximity to allow radiationless transfer of energy between the fluorochromes to form label complexes $(S_1\kappa D)_{n1}$ $(S_2\kappa A)_{n2}$ or $(S_1\kappa A)_{n1}$ $(S_2\kappa D)_{n2}$, wherein $n1$ and $n2$ represent the stoichiometry of the labeled subunits.

39. The method of claim 38, wherein S_1 is the regulatory subunit of cAMPdPK and S_2 is the catalytic subunit of cAMPdPK.

40. The method of claim 39, wherein S_2 is the regulatory subunit of cAMPdPK and S_1 is the catalytic subunit of cAMPdPK.

41. The method of claim 38, wherein the fluorochromes are fluorescein tetramethylrhodamine or derivatives thereof.

42. The method of claim 38, wherein the analyte is cAMP.

43. The method of claim 38, wherein S_1 and S_2 are recombinant proteins.

44. The method of claim 38, wherein one protein is a recombinant protein and one protein is derived from mammalian sources.

45. The method of claim 38, wherein S_1 and S_2 are derived from mammalian sources.

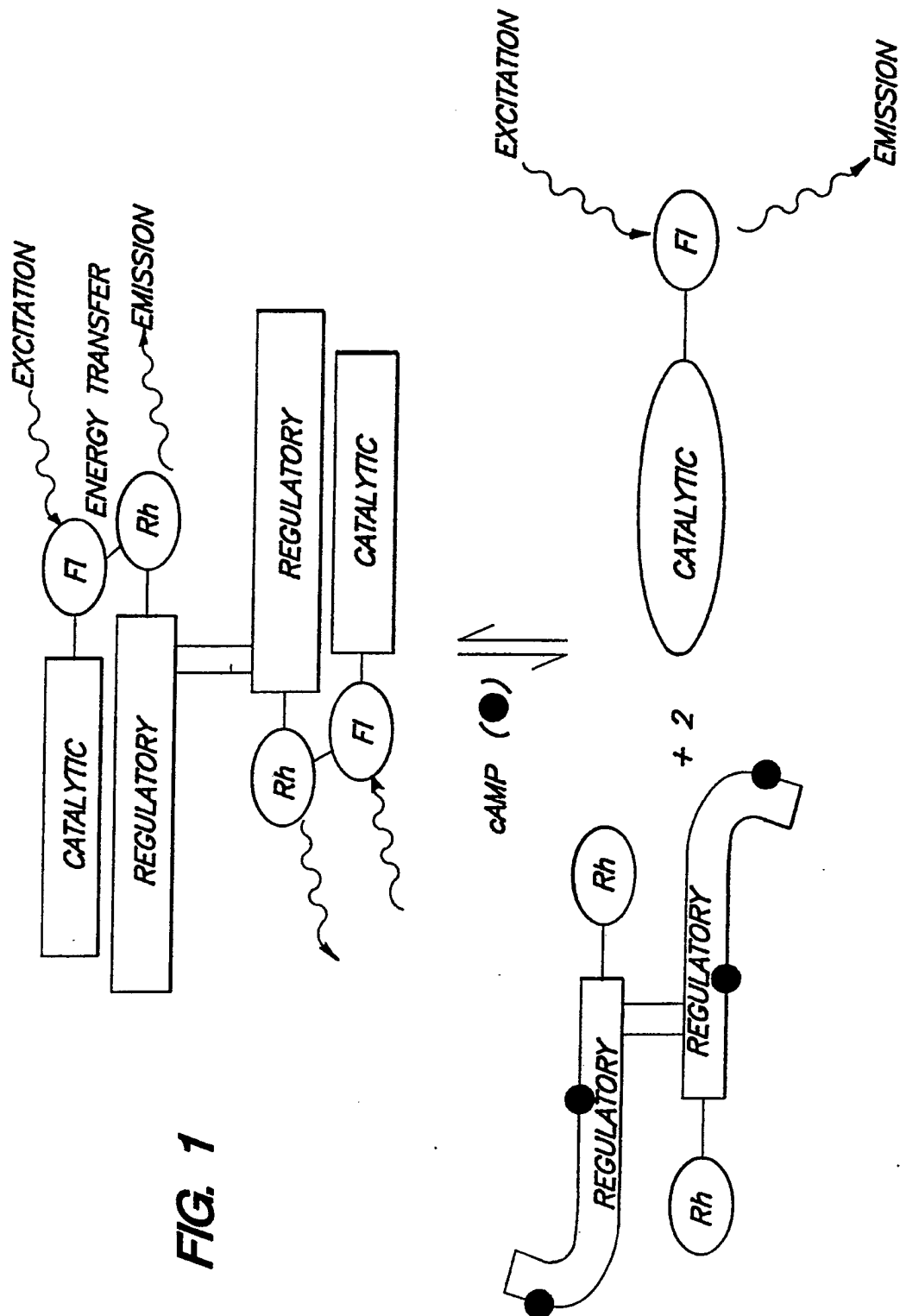


FIG. 1

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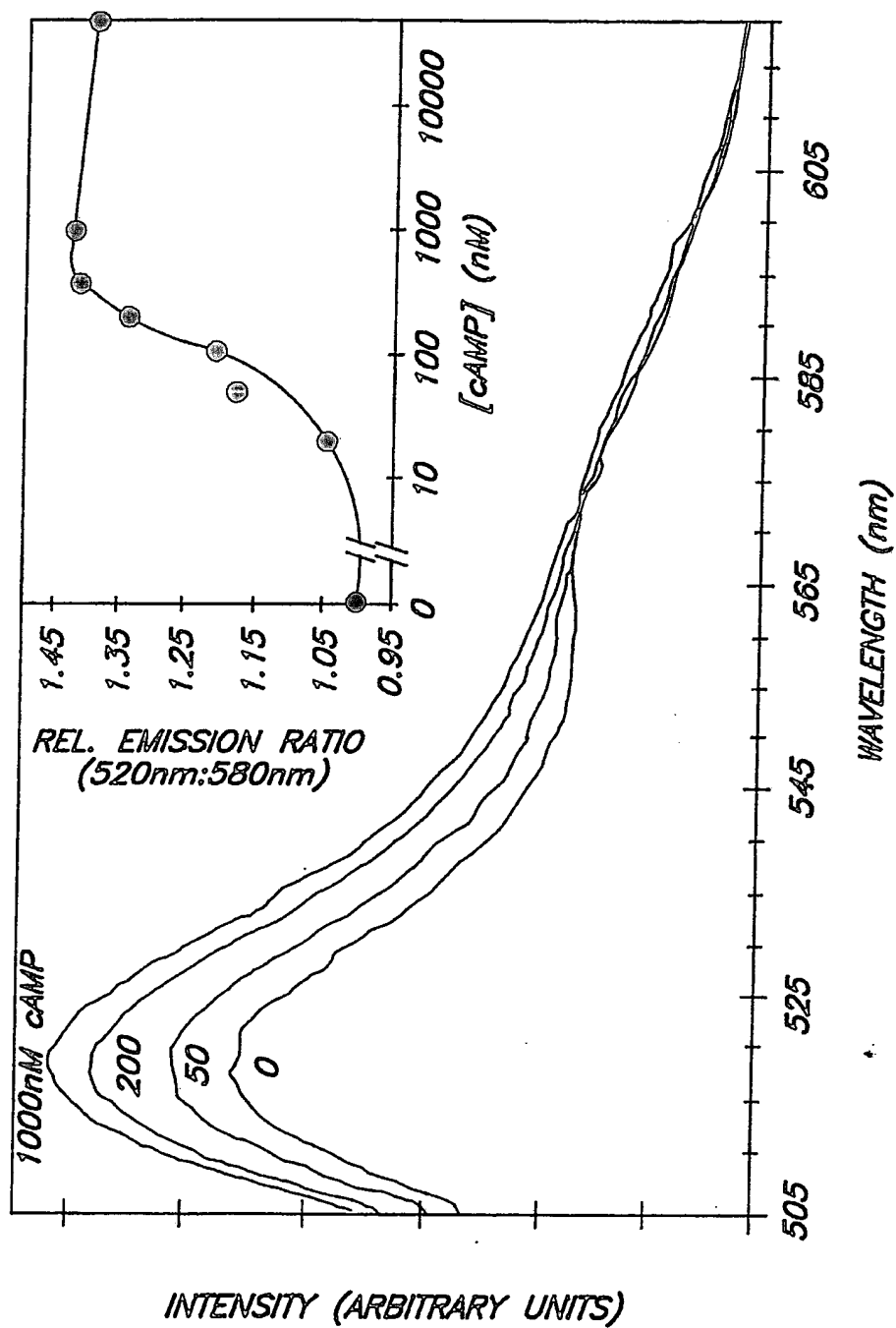
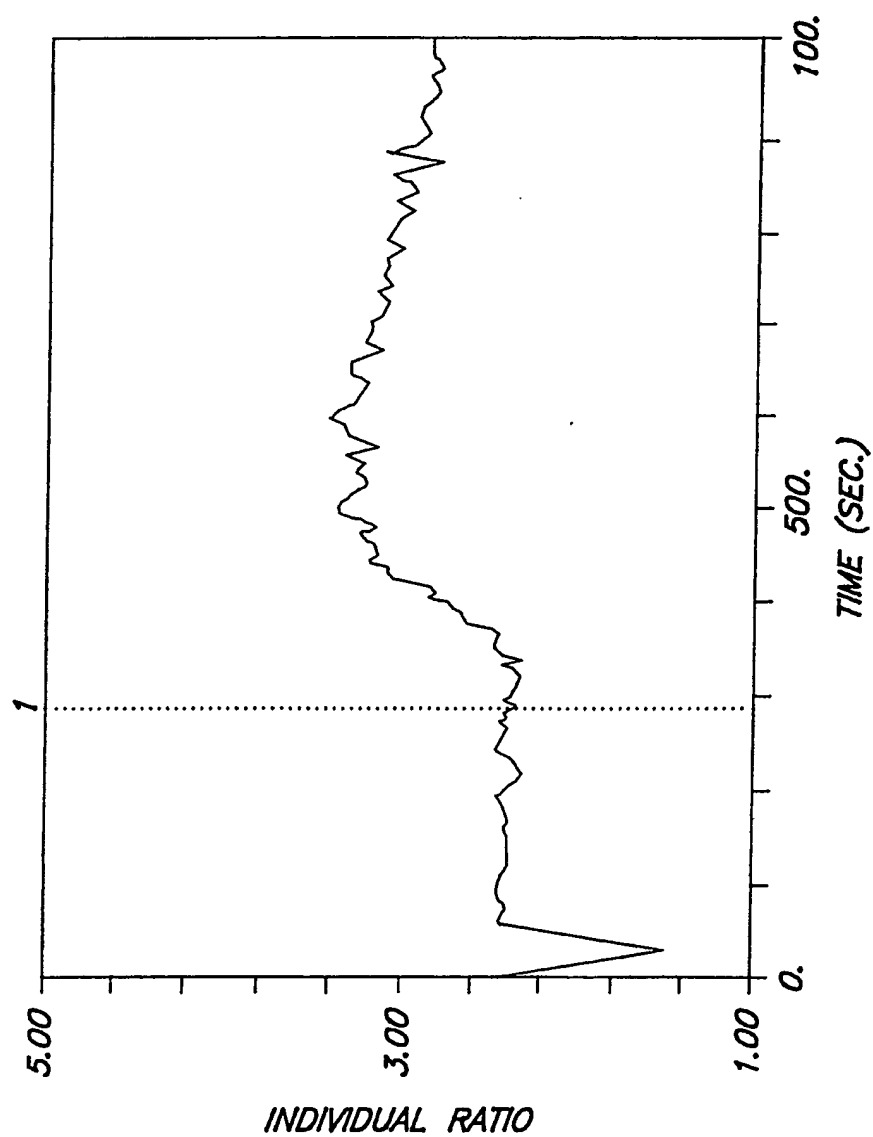


FIG. 2

SUBSTITUTE SHEET

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**FIG. 3****SUBSTITUTE SHEET**

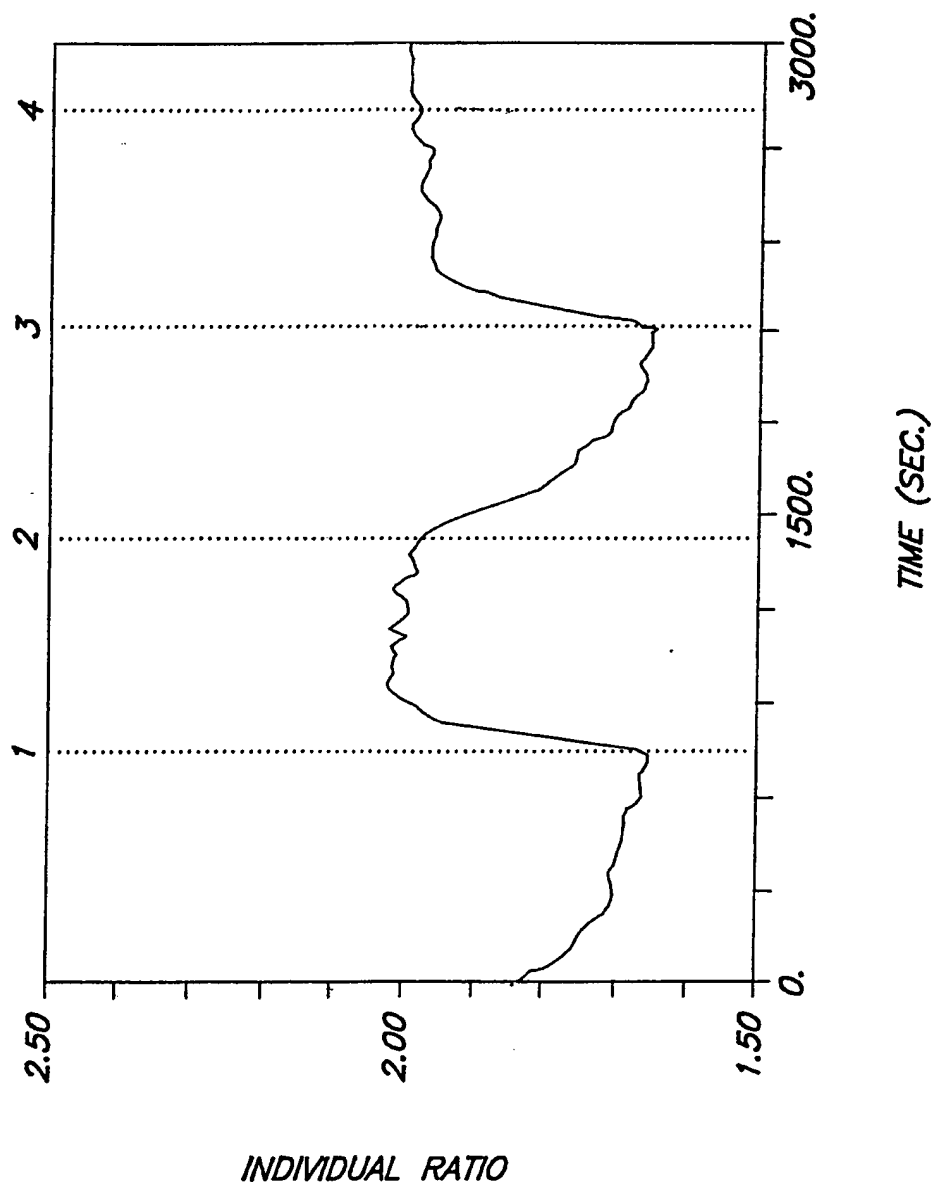


FIG. 4

INTERNATIONAL SEARCH REPORT

International Application No. PCT/US91/04676

I. CLASSIFICATION OF SUBJECT MATTER (If several classification symbols apply, indicate all) * According to International Patent Classification (IPC) or to both National Classification and IPC U.S. CL.: 435/7.21; 436/501, 546 IPC(5): C120 1/68; G01N 33/566, 33/533																	
II. FIELDS SEARCHED <div style="text-align: center; border-top: 1px solid black; border-bottom: 1px solid black;">Minimum Documentation Searched ?</div> <table style="width: 100%; border-collapse: collapse;"> <tr> <th style="width: 25%; border: 1px solid black;">Classification System</th> <th style="border: 1px solid black;">Classification Symbols</th> </tr> <tr> <td style="border: 1px solid black; vertical-align: top; padding: 5px;">U.S. CL.</td> <td style="border: 1px solid black; vertical-align: top; padding: 5px;">435/7.21; 436/501, 436/546</td> </tr> </table> <div style="text-align: center; border-top: 1px solid black; border-bottom: 1px solid black;">Documentation Searched other than Minimum Documentation to the Extent that such Documents are Included in the Fields Searched *</div> APS, MEDLINE SEARCH TERMS: energy transfer, CAMP dependent protein kinase, association, dissociation, subunit			Classification System	Classification Symbols	U.S. CL.	435/7.21; 436/501, 436/546											
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U.S. CL.	435/7.21; 436/501, 436/546																
III. DOCUMENTS CONSIDERED TO BE RELEVANT * <table border="1" style="width: 100%; border-collapse: collapse;"> <thead> <tr> <th style="width: 10%; border: 1px solid black;">Category *</th> <th style="border: 1px solid black;">Citation of Document, ** with Indication, where appropriate, of the relevant passages **</th> <th style="border: 1px solid black;">Relevant to Claim No. **</th> </tr> </thead> <tbody> <tr> <td style="border: 1px solid black; text-align: center; vertical-align: top;">X Y</td> <td style="border: 1px solid black; vertical-align: top;">US, A, 4,868,103, (STAVRIANOPOULOS ET AL.) 19 September 1989, see Col.5, lines 20-60.</td> <td style="border: 1px solid black; vertical-align: top;">1,2,5-7,14-16,19 22-24 & 31</td> </tr> <tr> <td style="border: 1px solid black; text-align: center; vertical-align: top;">X Y</td> <td style="border: 1px solid black; vertical-align: top;">US, A, 4,828,981, (MAGGIO) 09 May 1989, see entire document.</td> <td style="border: 1px solid black; vertical-align: top;">1-45 1,5-7,14-16,22-24 & 31</td> </tr> <tr> <td style="border: 1px solid black; text-align: center; vertical-align: top;">Y</td> <td style="border: 1px solid black; vertical-align: top;">J. BIOLOGICAL CHEMISTRY, Vol. 261, issued 26 April 1986, Fletcher et al., "Cytochemical identification of the regulatory subunit of the cAMP-dependent protein kinase by use of fluorescently labeled catalytic subunit," pages 5504-5513, see entire document.</td> <td style="border: 1px solid black; vertical-align: top;">1-45 17,18</td> </tr> <tr> <td style="border: 1px solid black; text-align: center; vertical-align: top;">Y</td> <td style="border: 1px solid black; vertical-align: top;">J. BIOLOGICAL CHEMISTRY, Vol. 264, issued 05 August 1989, Woodford et al., "Expression and characterization of mutant forms of the type I regulatory subunit of cAMP-dependent protein kinase," pages 13321-13328, see entire document.</td> <td style="border: 1px solid black; vertical-align: top;">11-13,28-30,35-37 43-45</td> </tr> </tbody> </table>			Category *	Citation of Document, ** with Indication, where appropriate, of the relevant passages **	Relevant to Claim No. **	X Y	US, A, 4,868,103, (STAVRIANOPOULOS ET AL.) 19 September 1989, see Col.5, lines 20-60.	1,2,5-7,14-16,19 22-24 & 31	X Y	US, A, 4,828,981, (MAGGIO) 09 May 1989, see entire document.	1-45 1,5-7,14-16,22-24 & 31	Y	J. BIOLOGICAL CHEMISTRY, Vol. 261, issued 26 April 1986, Fletcher et al., "Cytochemical identification of the regulatory subunit of the cAMP-dependent protein kinase by use of fluorescently labeled catalytic subunit," pages 5504-5513, see entire document.	1-45 17,18	Y	J. BIOLOGICAL CHEMISTRY, Vol. 264, issued 05 August 1989, Woodford et al., "Expression and characterization of mutant forms of the type I regulatory subunit of cAMP-dependent protein kinase," pages 13321-13328, see entire document.	11-13,28-30,35-37 43-45
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<div style="display: flex; justify-content: space-between;"> <div style="width: 45%;"> <p>* Special categories of cited documents: **</p> <p>"A" document defining the general state of the art which is not considered to be of particular relevance</p> <p>"E" earlier document but published on or after the international filing date</p> <p>"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)</p> <p>"O" document referring to an oral disclosure, use, exhibition or other means</p> <p>"P" document published prior to the international filing date but later than the priority date claimed</p> </div> <div style="width: 45%;"> <p>"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention</p> <p>"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step</p> <p>"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art.</p> <p>"&" document member of the same patent family</p> </div> </div>																	
IV. CERTIFICATION <table style="width: 100%; border-collapse: collapse;"> <tr> <td style="width: 50%; border: 1px solid black; vertical-align: top; padding: 5px;"> Date of the Actual Completion of the International Search 30 SEPTEMBER 1991 </td> <td style="width: 50%; border: 1px solid black; vertical-align: top; padding: 5px;"> Date of Mailing of this International Search Report 29 NOV 1991 </td> </tr> <tr> <td style="border: 1px solid black; vertical-align: top; padding: 5px;"> International Searching Authority ISA/US </td> <td style="border: 1px solid black; vertical-align: top; padding: 5px;"> Signature of Authorized Officer M.P. WOODWARD </td> </tr> </table>			Date of the Actual Completion of the International Search 30 SEPTEMBER 1991	Date of Mailing of this International Search Report 29 NOV 1991	International Searching Authority ISA/US	Signature of Authorized Officer M.P. WOODWARD											
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III. D CUMENTS CONSIDERED TO BE RELEVANT (C NTINUED FROM THE SECOND SHEET)

Category *	Citation of Document, with indication, where appropriate, of the relevant passages	Relevant to Claim No
X Y	J. BIOLOGICAL CHEMISTRY, Vol. 261, issued 25 February 1986, McEvily et al., "Subunit equilibria of porcine heart citrate synthase," pages 2593-2598, see entire document.	1,2,5-7,14-16,19,22-24 & 31 1-7,12-16,19,22-24 30,31,33,37,38,41& 45
Y	BIOCHEMISTRY, Vol. 28, issued 1989, First et al., "Fluorescence energy transfer between cysteine 199 and cysteine 343: evidence for MgATP-dependent conformational change in the catalytic subunit of cAMP-dependent protein kinase," pages 3606-3613, see entire document.	1-45
Y	BIOCHEMISTRY, Vol. 28, issued 1989, First et al., "Selective modification of the catalytic subunit of cAMP-dependent protein kinase with sulfhydryl-specific fluorescent probes," pages 3598-3605, see entire document.	1-45
Y	EUR. J. BIOCHEM., Vol. 188, issued 1990, Buchler et al., "Regulation of gene expression by transfected subunits of cAMP-dependent protein kinase," pages 253-259, see entire document.	11-13,17,18,28-30 35-37&43-45